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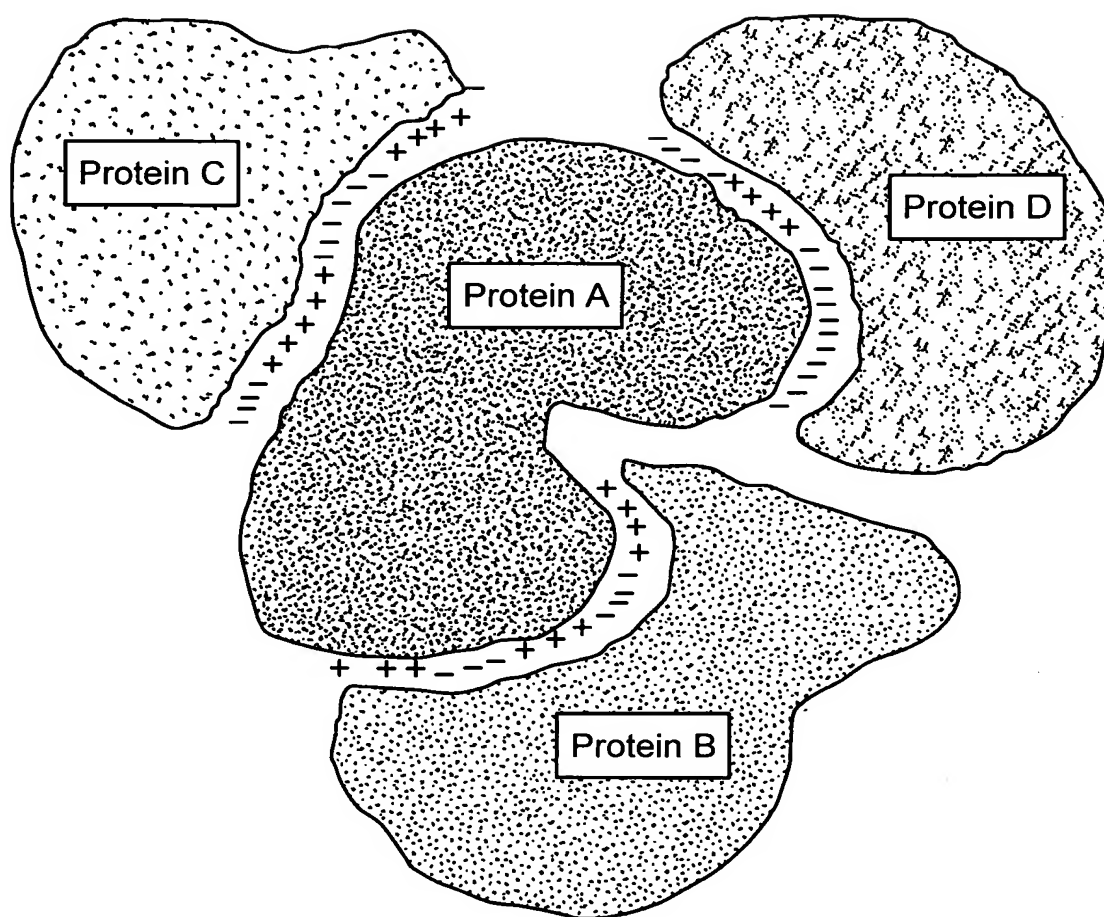


FIG. 1

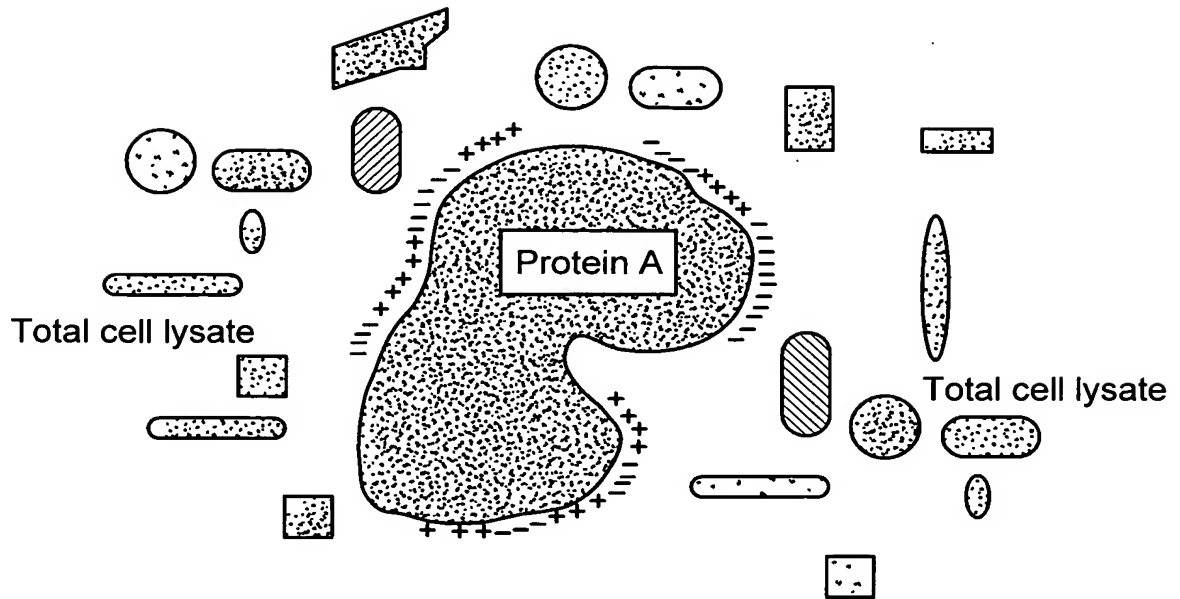


FIG. 2A

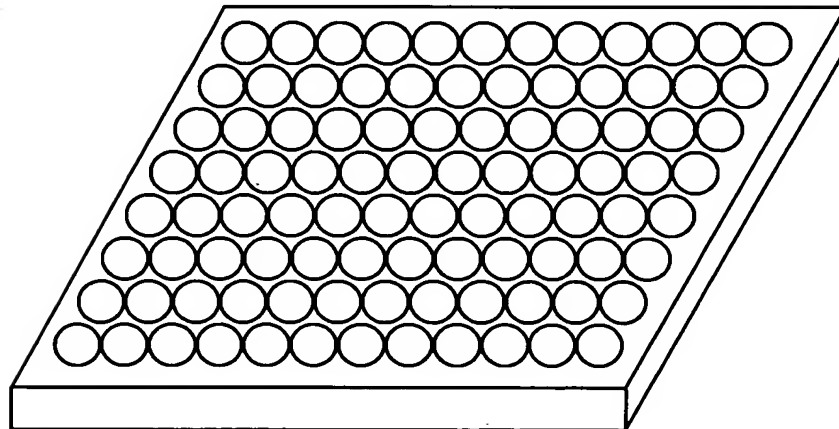


FIG. 2B



3/16

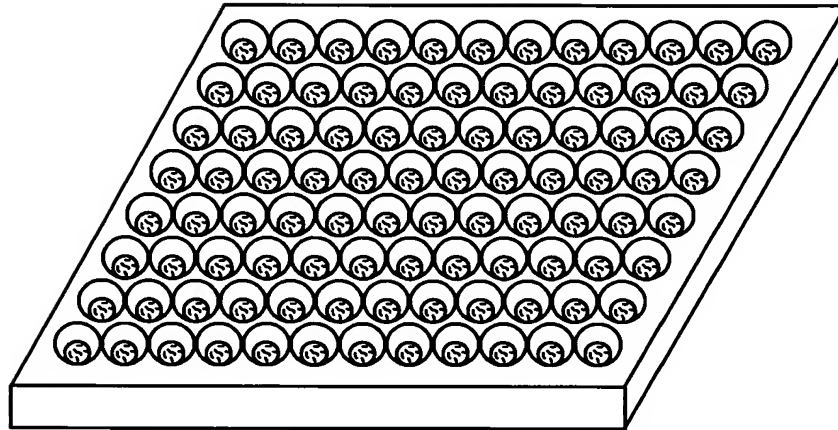


FIG. 2C

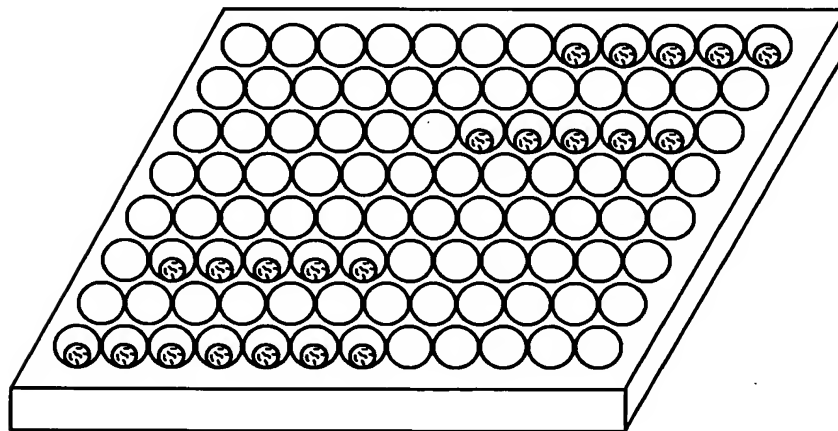


FIG. 2D

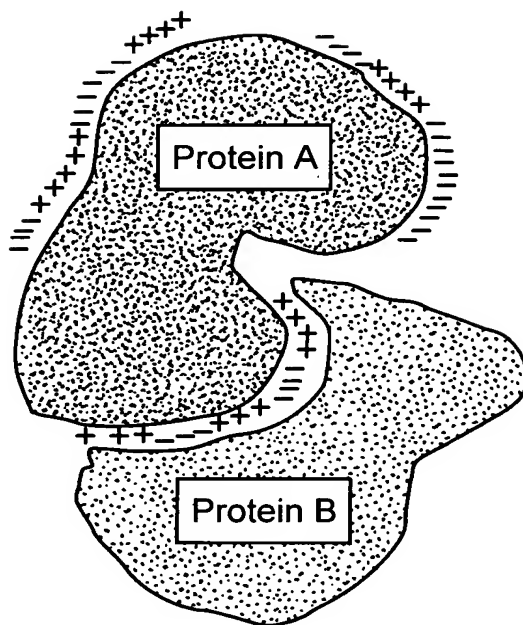


FIG. 3A

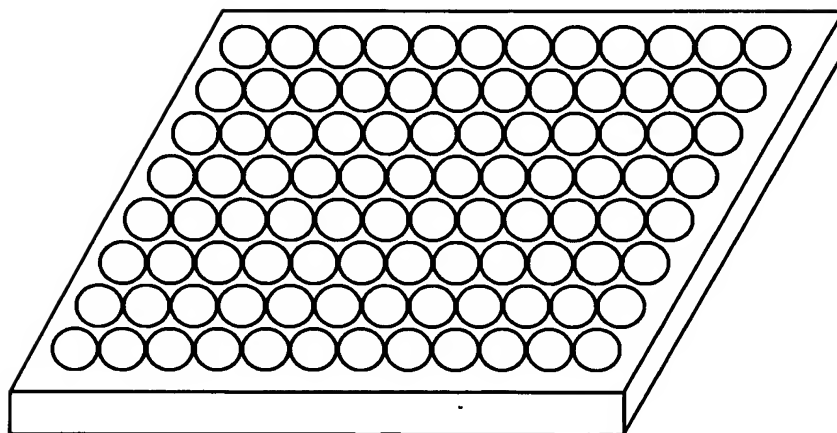


FIG. 3B

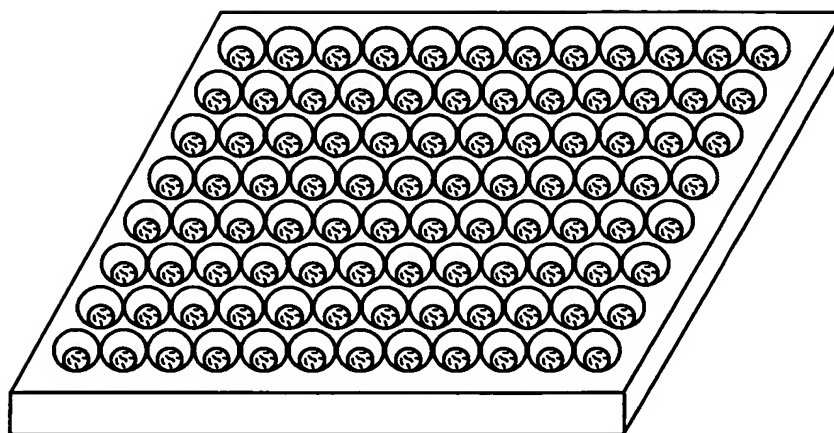


FIG. 3C

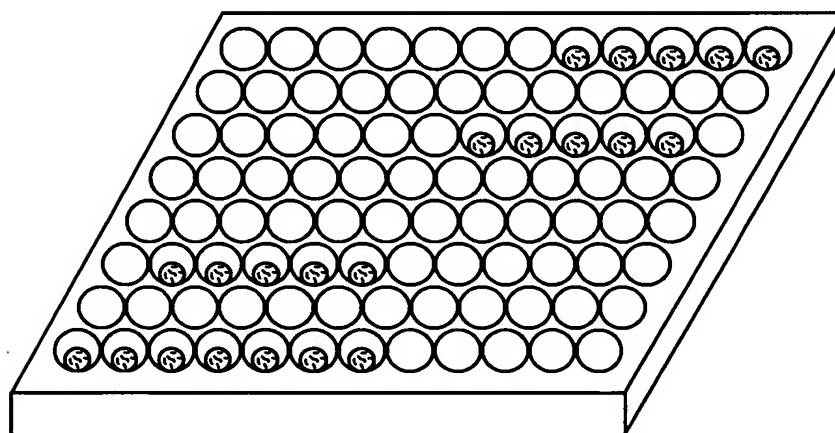


FIG. 3D

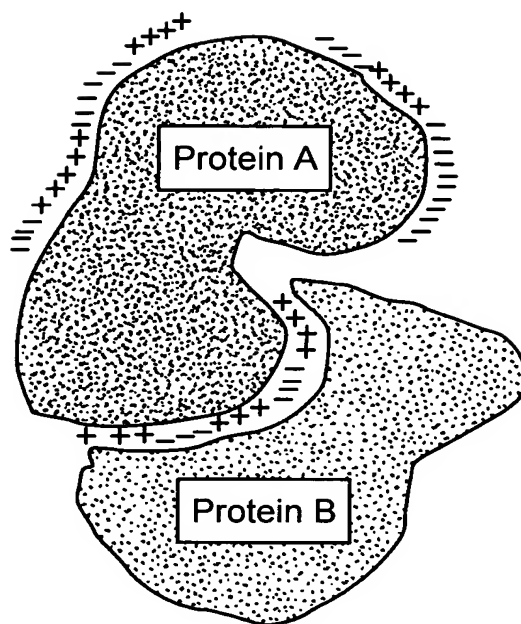


FIG. 4A

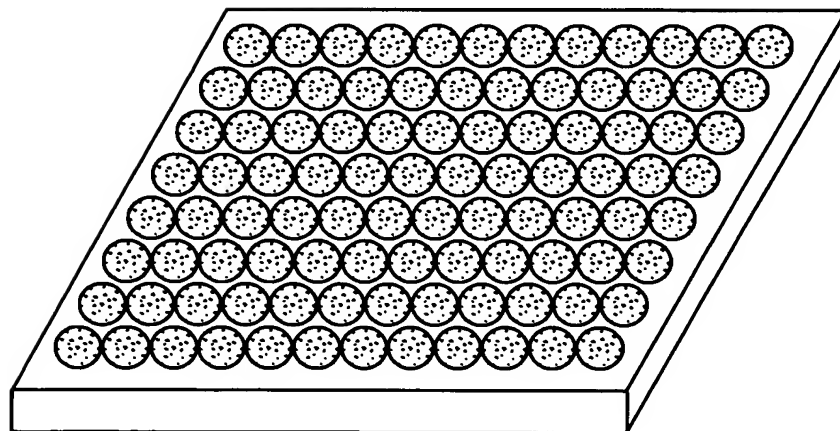


FIG. 4B

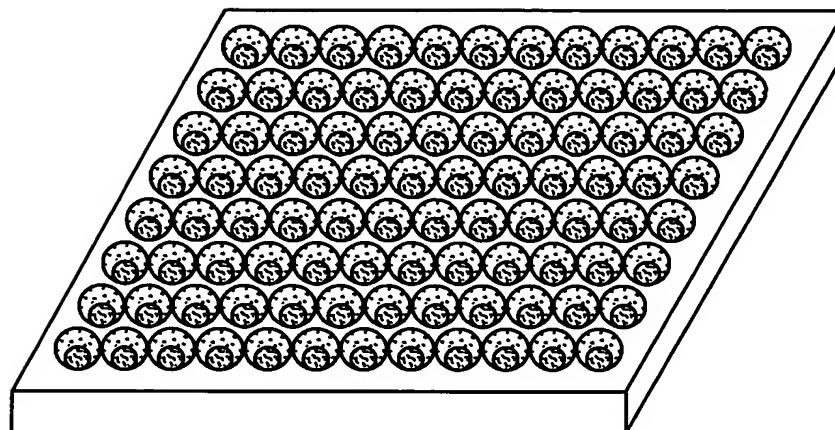


FIG. 4C

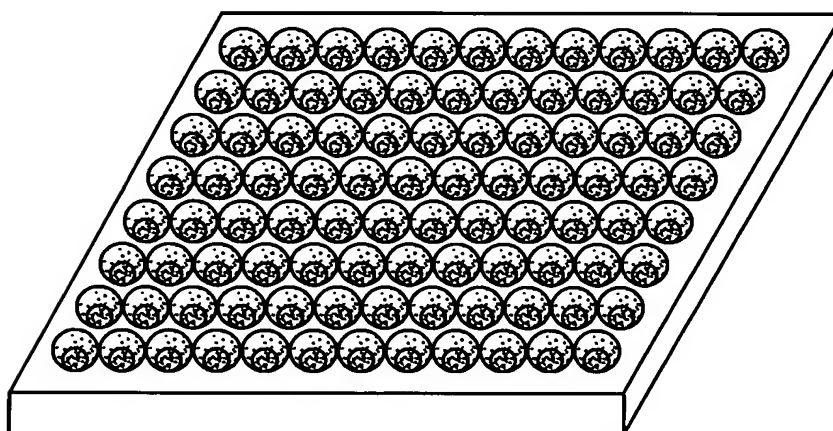


FIG. 4D

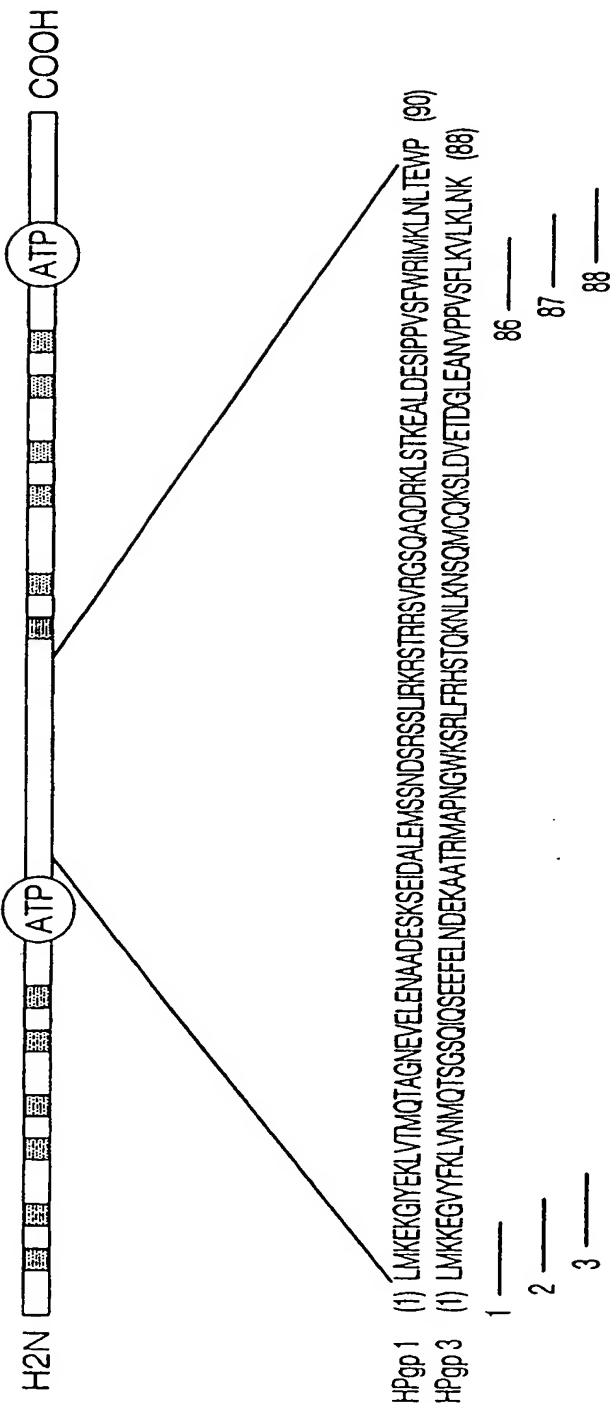
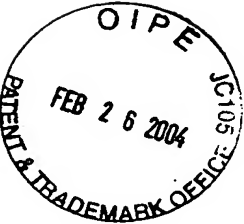


FIG. 5

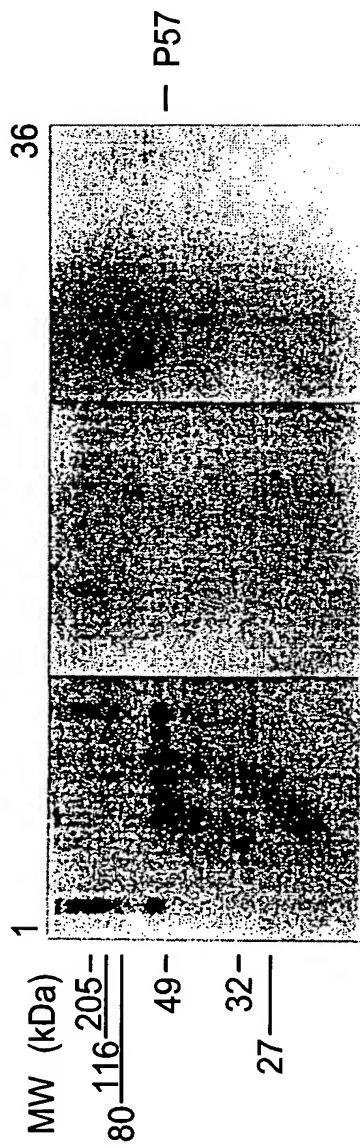


FIG. 6A

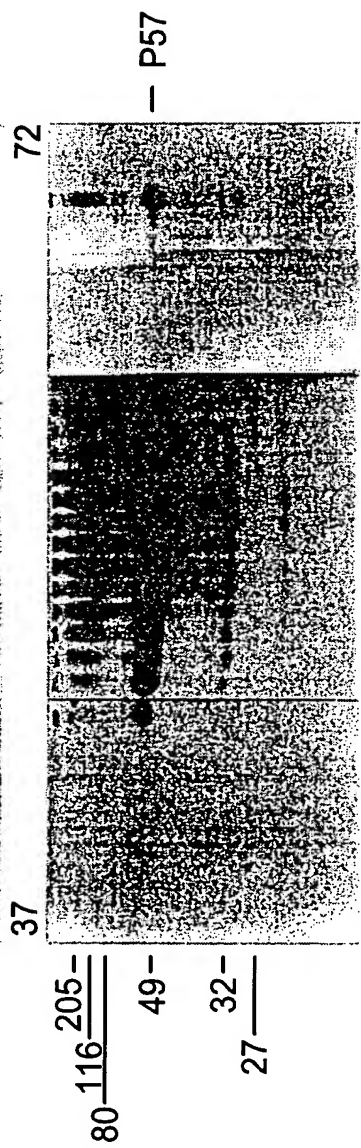


FIG. 6B

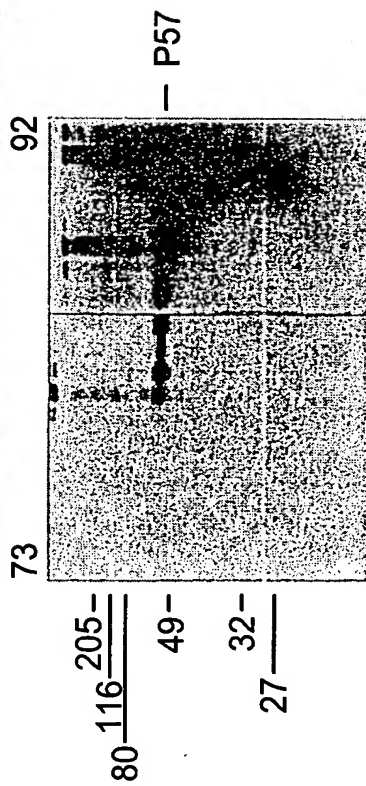


FIG. 6C

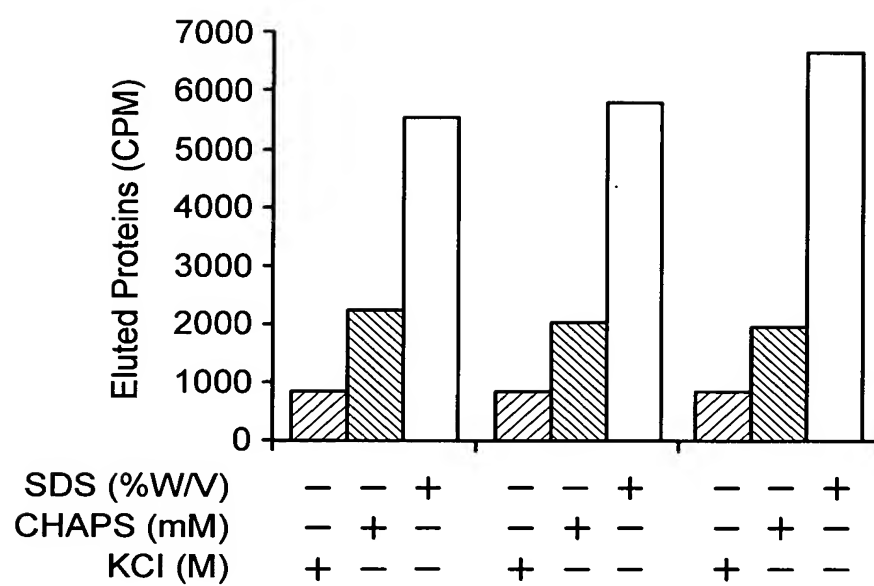


FIG. 7



11/16

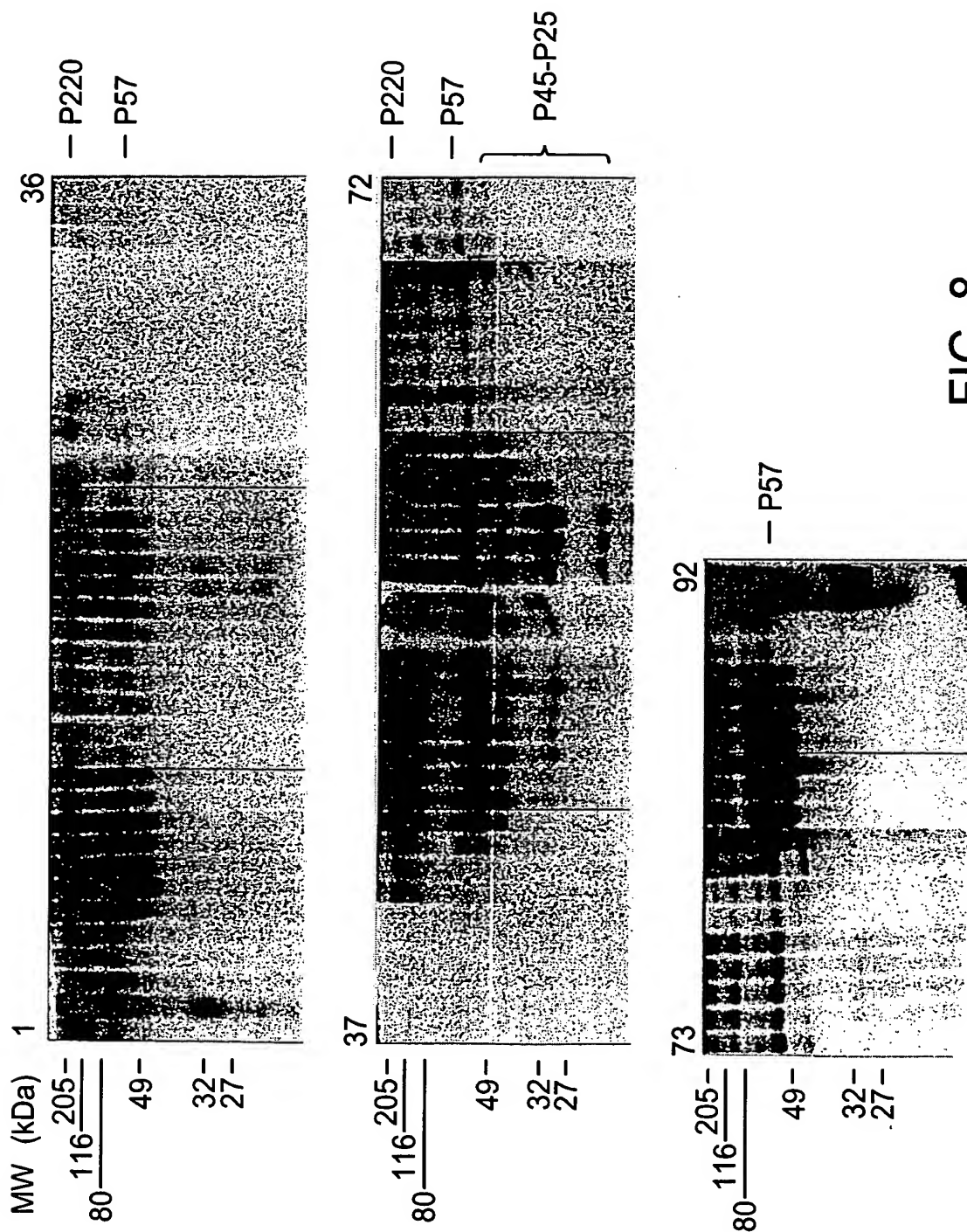


FIG. 8



12/16

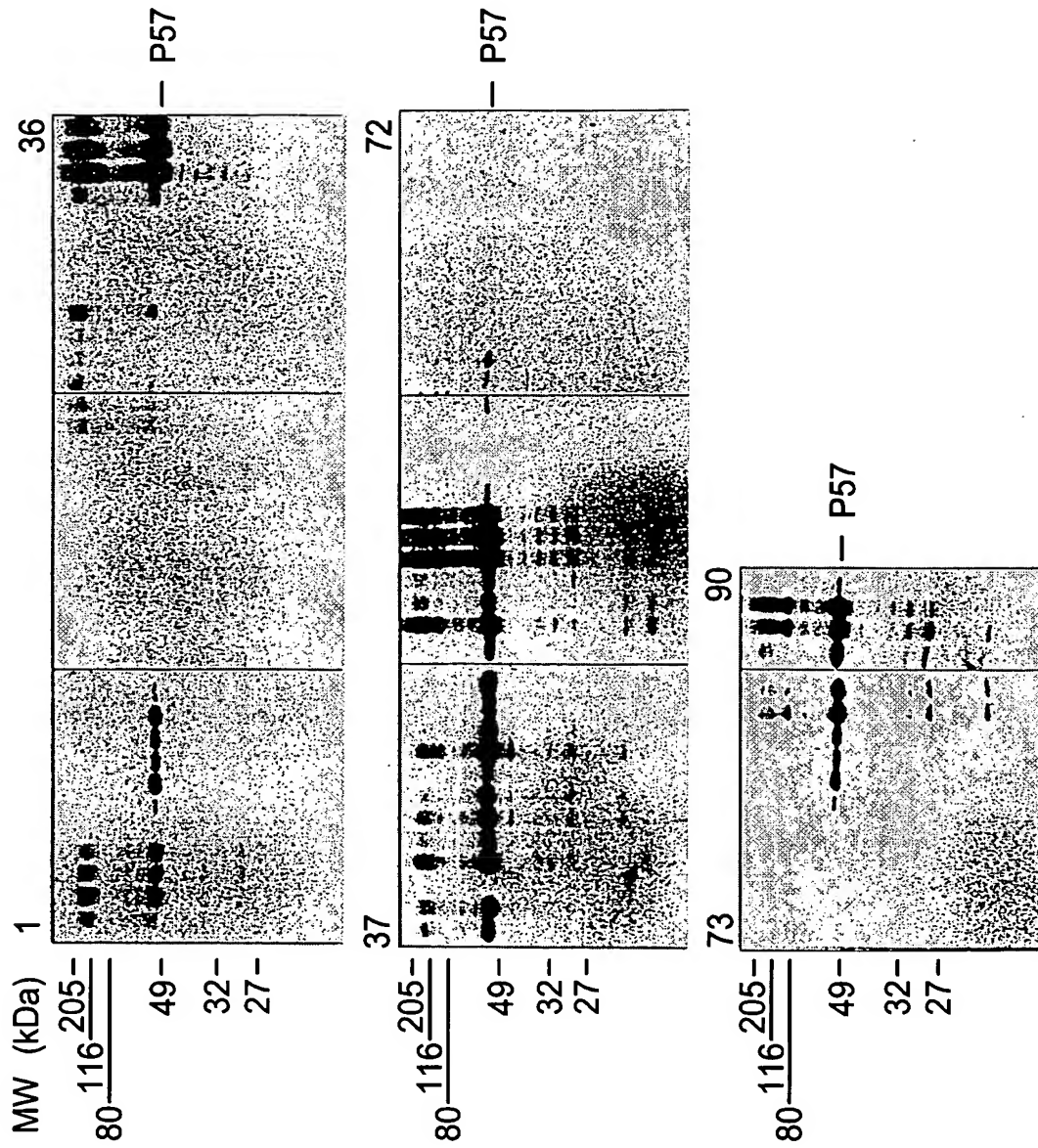
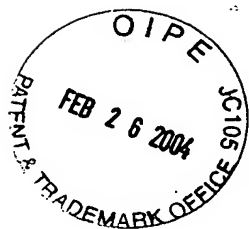


FIG. 9



Alignment Of Amino Acid Sequences Of Human P-gp3 And P-gp1
Linker Domains

618 LMKKEGYFKLVNMQTSGSQIQSEEF--ELNDEKAATRMAPNGHKSRLFR-HSTQKNLKN SQM 677 P-gp3

LMK++G+YFKLV MQT+G+++ E E E A M+ N +S L R ST++++

615 LMKKEGYFKLVMTMQTAGNEVELENAADESKSEIDALEMSSNDSRSLIRKSTRRSVRSQA 677 P-gp1

678 CQKSLDVETDGLANVPPVSFLKVLKLNKTEWP 710 P-gp3

+ L + + L+ ++PPVSF +++KLN TEWP

678 QDRKLSTK-EALDESIPPVSFWRIMKLNLTWP 709 P-gp1

FIG. 10

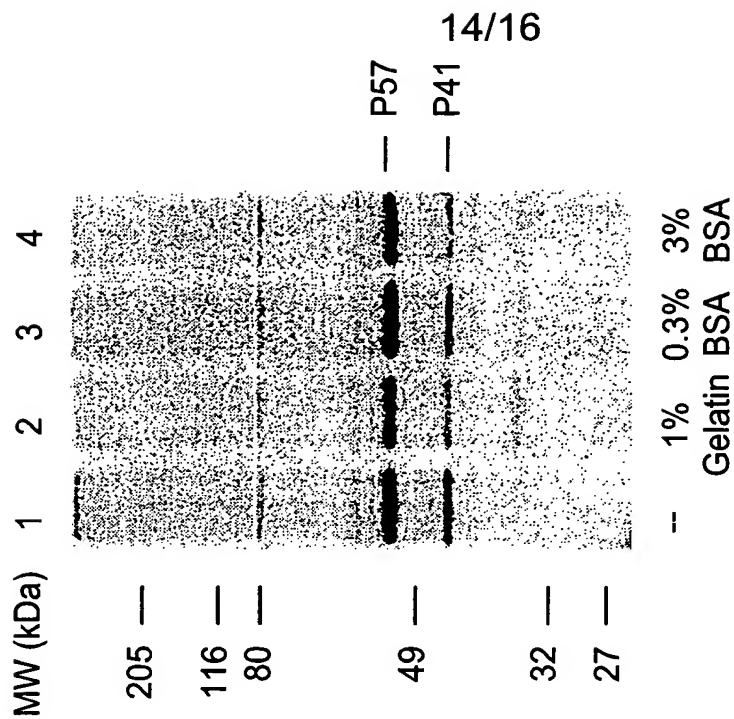
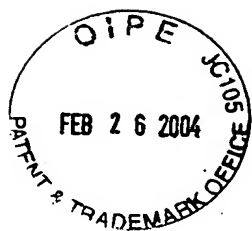


FIG. 12

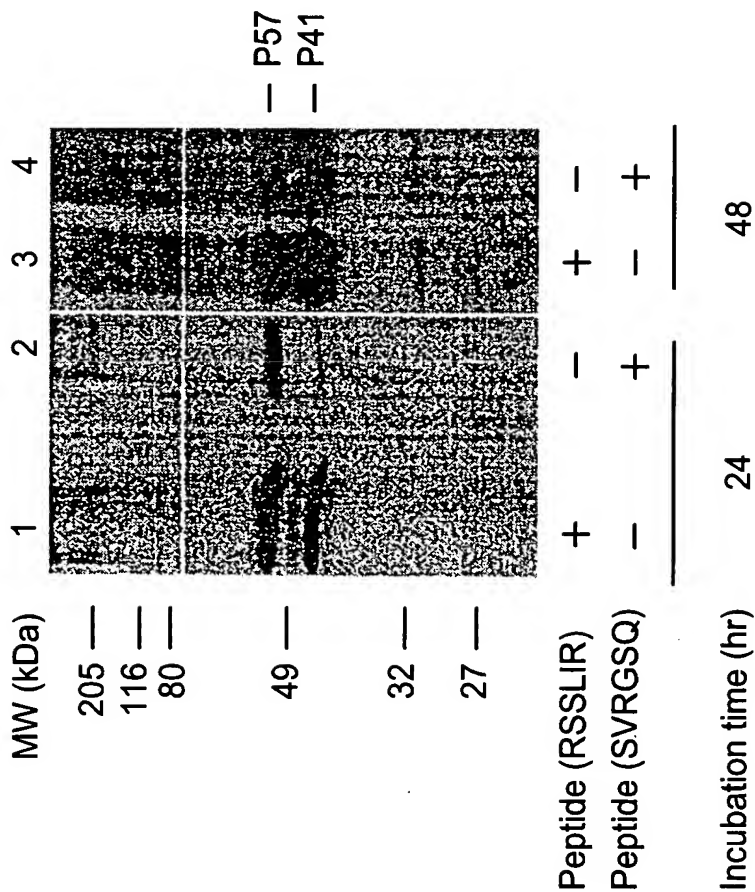


FIG. 11

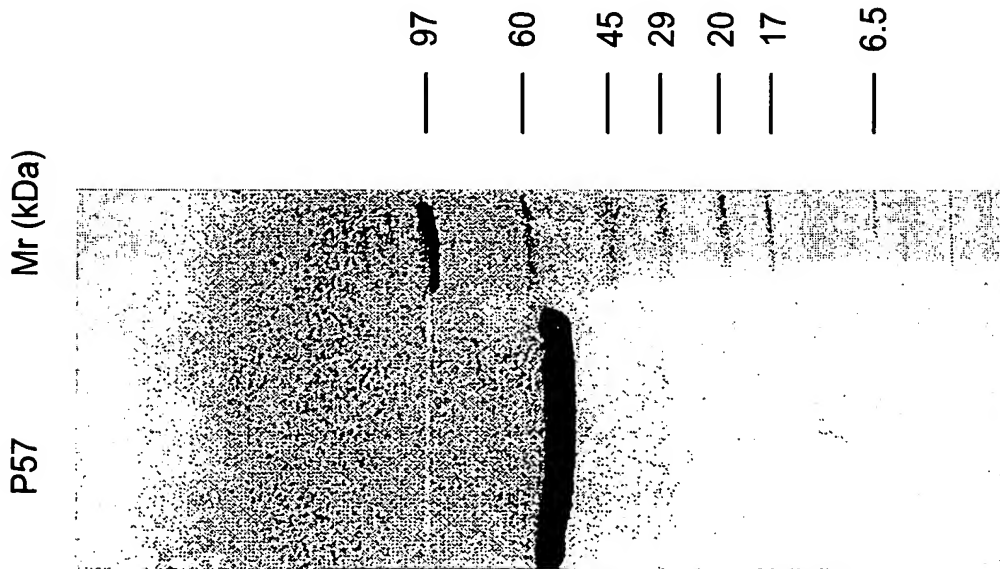
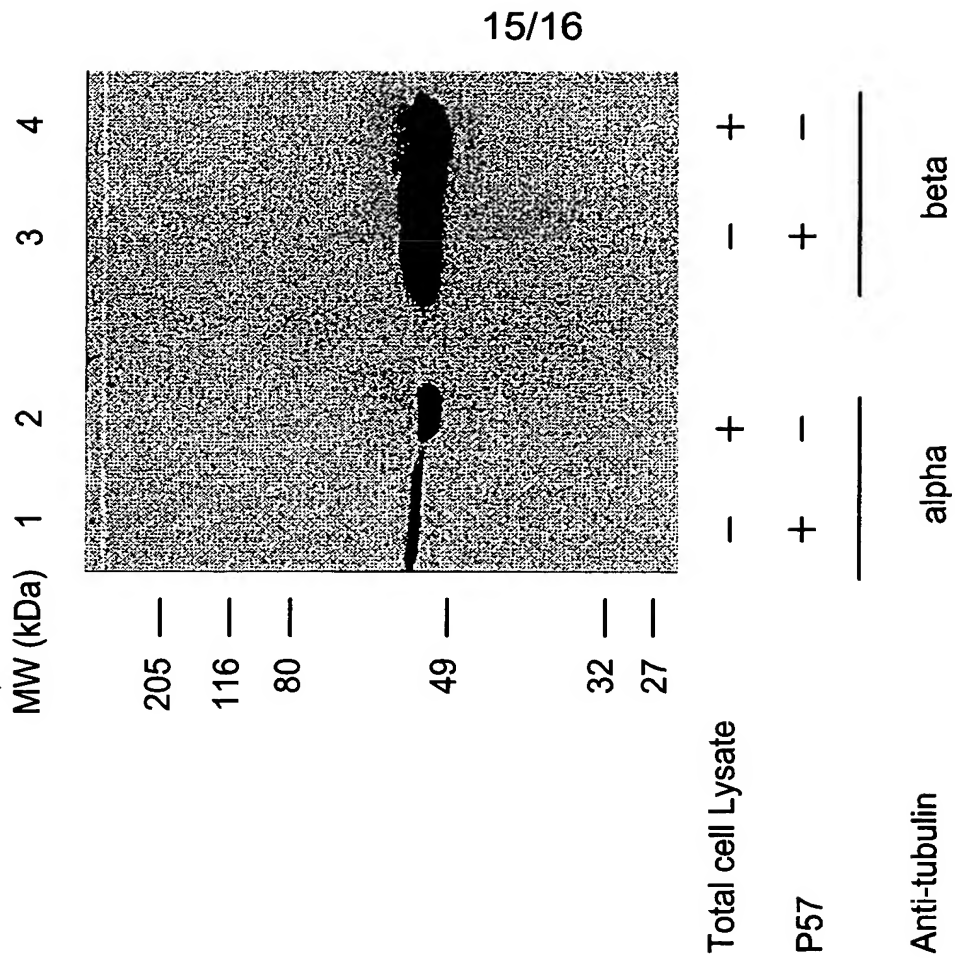


FIG. 13



Total cell Lysate
P57
Anti-tubulin

-	+	-	+
+	-	+	-
alpha		beta	

FIG. 14



HELICAL WHEEL PRESENTATION

SRSSLIRKRSTRRSVRGS

NGWKSRLFRHSTQKNLK

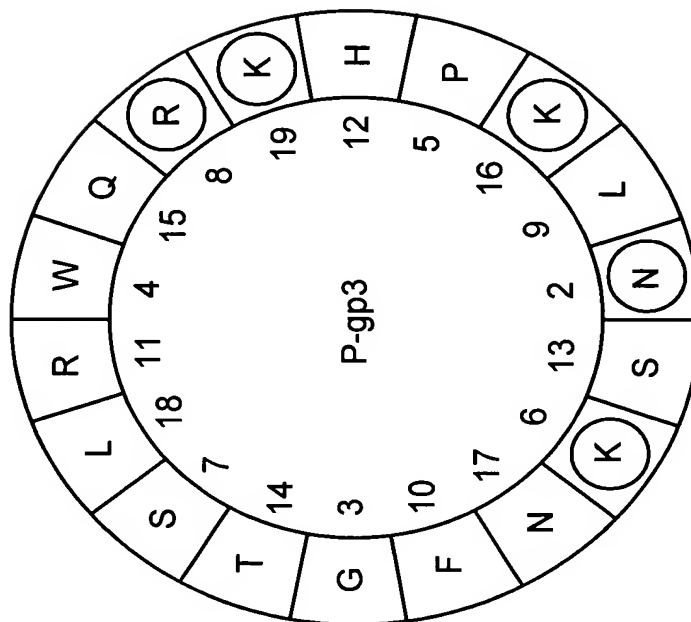
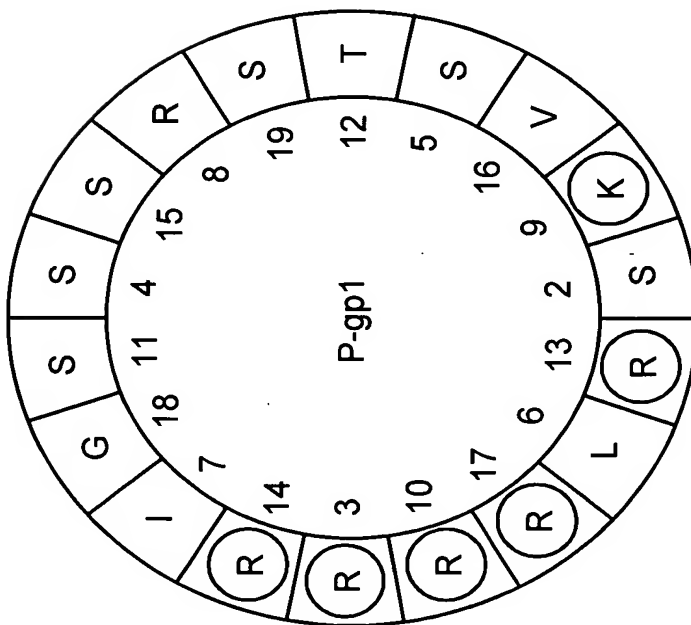


FIG. 15